# THE PERSONAL GENOME 

## Everyone has one

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## BIOINFORMATIC CONSIDERATIONS

## MASSIVE AMOUNTS OF DATA



EACH GENOME GENERATES 100 GB DATA FOR DOWNSTREAM ANALYSIS, EACH GENOME GENERATES 100 GB DATA FOR DOWNSTREAM ANALYSIS,
THIS REOUIRES STORAGE BEYOND WHAT THE TYPICAL HPC ACCOMMODAT 100 MILLION GENOMES X 100 GIGABYTES OF DATA $=10$ EXABYTES OF DATA

> NEED FOR INNOVATIVE TECHNOLOGY $\&$ CLOUD-NATIVE ANALYSIS

- ngS TOOLS ARE NOT CROSS-PLATFORM TOOLS. THE MAJORITY OF THEM WORK O the linux/unix operating system. -MOST OF THE NGS TOOLS INTERACT WITH
USERS VIA COMMAND LINE INTERFACE.
- VERY few ngs tools have been解 THERE IS NO STANDARD INPUT/OUTPUT FILE
FORMAT FO

RESULT: REDUCED COMPATIBILITY, PORTABILITY,
INTEROPERABILITY, AND INTEGRATION BETWEEN THE TOOLS.
FINDINGS FROM BANI BAKER ET AL. 2020 (DOI.ORG/10.1016/J.IMU.2020.100296)

## SECURITY IN DATA COLLABORATIONS

"RESEARCHERS MUST bE ABLE TO RESPONSIBLY INTERROGATE THE WORLD'S COLLECTIVE GENOMIC DATA
AS A SINGLE VIRTUAL COHORT THAT TRANSCENDS THE CAPACITY FROMD JURISDICTION - OF ANY SINGLE ORGANISATION OR COUNTRY. FROM ARTICLE - "ELIXIR AND GA4GH EXPAND COLLABORATION", MAY 2019


STARK ET AL. NTEGRATING GENOMICS INTO HEALTHCARE: A GLOBAL RESPONSIBLITY.
THE AMERICAN JOURNAL OF HUMAN GENETICS 104.1 (2019): 13 -20.

## NEED FOR CLOUD SOLUTIONS

PERSONAL GENOMES CAN BE COMPRESSED AND CONVERTED TO SEARCHABLE FORMATS

CLOUD SIMPLIFIES MAKING DATA F.A.I.R. = FINDABLE ACCESSIBLE, INTEROPERABLE AND REUSABLE

- CONVERT GENOMIC vCF INTO COMPRESSED BIGQUERY OR ATHENA - CORMERTS
FORMATS
- USE REST API AND SQL TO SEARCH VARIANTS ACROSS GENOMES WITH -USE REST API AND SQL TO SEARCH VARIANTS A
SIMPLE COORDINATES OR PATTERN MATCHING
- multiple variant sets for pangenomes can be joined with simple SQL COMMANDS
- PERSONAL genome data can be link to public test data sets such as
1000 genomes, simon genomes, platinum genomes.
- RAW SEqUENCING DATA CAN be moved to archival storage once
ALIGNMENTS ARE PRODUCED.

PERSONAL GENOMES BENEFIT FROM
SCALABLE OPTIONS FOR
MACHINE LEARNING TO OPTIMIZE COSTS
AND REDUCE TIME

## KUBEFLOW

- UTLIZES GPU'S (GRAPHICAL PROCESSING UNITS) TO SPEED UP processing
- PARTITION MULTIPLE JOBS ON ONE GPU
- CAN be controlled from local machine or from a vm in gcp TENSORFLOW
- ALLOWS EXPANSION OF ML to tpu's (tensor processing units) - higher memory options now available
- takes minutes to complete versus days or weeks.

PERSONAL GENOMES IN THE CLOUD CAN HAVE AUTOMATED ENCRYPTION AND SECURITY POLICIES
develop and automate compliance \& cyber risk management PROCESSES

- COMPLY wITH FISM, NIST RMF AND HIPAA
- SECURE GENOMICS DATA, PHI AND CLINICAL\TRANSLATIONAL
- SECURE GENOMICS DATA, PHI AND CLINICAL
RESEARCH DATA IN CLOUD ENVIRONMENTS
- AUTOMATE GA4GH, SSO AND CLOUD USAGE POLICIES

ONIX ENABLES HEALTH AND LIFE SCIENCE ORGANIZATIONS to realize the power of the cloud
TO SECURELY WORK SIMPLER, SMARTER AND FASTER IN A HIPAA-COMPLIANT ENVIRONMENT.

GASPAR, H.A., BREEN, $\quad$. PROBABLLISTIC ANCESTRY MAPS: A
METHOD TO ASESES AND VISUALIZE POPULATION SUBSTRUCTURES IN GENETICS. BMC BIOINFORMATICS 20, 116 (2019).
HTTPS:/DOI.ORG/10.1186/S12859-019-2680-I

